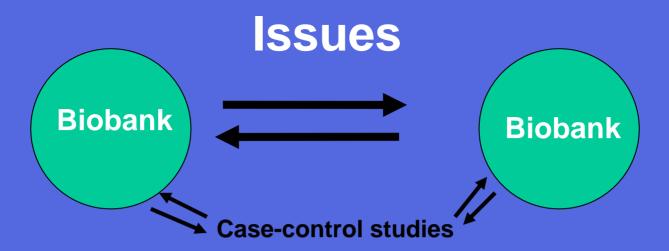


Professor Bill Ollier





- Observations and findings from one Biobank study need to be replicated and confirmed in others
- Small effects observed in one study may require a metaanalysis of multiple datasets for confirmation and precise estimation of size
- Hypotheses generated in large prospective population studies may need to be tested in the context of case-control studies and vice-versa
- Observations need to be tested and interpreted in a range of ethnic backgrounds
- Gene-environment interactions are likely to require metaanalysis

Consequences

These issues will require:

- Major national and international collaboration between large population projects in genetics with agreed terms of reference
- Agreement on what are minimum critical datasets and quality standards
- Introduction of harmonisation/standardisation and for phenotype/genotype definition and data collection and management

- Minimum standards for biological sample storage
- Comparability of environmental exposure data
- Compatibility of environmental exposure data
- Compatibility of statistical and analytical approaches
- Minimum descriptors for study design, analysis and reporting of results

Results

- Web-based detailed description of study design and protocol (Perhaps a centralised inventory?)
- Context in which dataset (subset) has been selected and defined e.g. disease phenotype definition and minimum critical data fields e.g. gender, age at disease onset, ethnicity
- On-line access to original datasets on which analysis has been performed, intermediate summary tables and results reported
- Predicted effect sizes
- Extent of multiple testing

Concerns

- Standardisation of procedures and reporting should not inhibit or stifle innovation of analytical approach, biological interpretation or hypothesis generation
- Confidentiality or ethical framework of original studies should not be compromised
- Secondary or meta-analysis of datasets may reveal new observations and interactions for which original investigators will not be acknowledged or IP credited